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Total number of hits satisfying chosen parameters:
                                                                                                                                                                    Title:
Perfect score:
                                                Searched:
                                                                                                       Scoring table:
                                                                                                                                                 Sequence:
                                                                                                                                                                                                                                                                       Run on:
                                                                                                                                                                                                                                                                                                            OM protein - protein search, using sw model
                                                                                    BLOSUM62
Gapop 10.0 , Gapext 0.5
                                            3502263 seqs, 351980561 residues
                                                                                                                                                                  US-09-502-984B-37
1284
                                                                                                                                               1 KFESKAALLAARGPEELLCF.....RKNERLEEEVERLKQLVGER 249
                                                                                                                                                                                                                                                  August 28, 2002, 17:37:38; Search time 305.46 Seconds (without alignments)
                                                                                                                                                                                                                                                                                                                                                                     GenCore version 4.5
Copyright (c) 1993 - 2000 Comp
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                                                                                                                                                                                                                          286.922 Million cell updates/sec
      3502263
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 10%
Maximum Match 100%
Listing first 45 summaries

Database: Pending_Patents_A_Main:*

1: /cgn2_6/ptodata/2/paa/US06_COMB.pep:*
2: /cgn2_6/ptodata/2/paa/US06_COMB.pep:*
3: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
4: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
5: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
6: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
8: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
9: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
9: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
11: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
11: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
11: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
12: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
13: /cgn2_6/ptodata/2/paa/US08_COMB.pep:*
14: /cgn2_6/ptodata/2/paa/US09_COMB.pep:*
15: /cgn2_6/ptodata/2/paa/US091_COMB.pep:*
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18: /cgn2_6/ptodata/2/paa/US093_COMB.pep:*
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23: /cgn2_6/ptodata/2/paa/US093_COMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/cgn2_6/ptodata/2/paa/US099_COMB.pep:*
/cgn2_6/ptodata/2/paa/US100_COMB.pep:*
/cgn2_6/ptodata/2/paa/US101_COMB.pep:*
/cgn2_6/ptodata/2/paa/US60_COMB.pep:*

SUMMARIES

765 4 321	Result No.
1171 1098 1080 1078 1075 1073 1073	Score
	Query Match 1
676 211 211 211 211 211 211 211	Length I
18 19 19 19 19	DB
US-09-452-565-1 US-09-502-984-6 US-09-502-984-5 US-09-502-984-4 US-09-502-984-9 US-09-502-984-13 US-09-502-984-14	ID
Sequence 1, Appli Sequence 6, Appli Sequence 5, Appli Sequence 4, Appli Sequence 9, Appli Sequence 13, Appl Sequence 14, Appl	Description

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3, A	Sequence 710, App	1197,	Sequence 2419, Ap	4200,	3517,	13860,	29, Ap	27,	Sequence 28, Appl	26,	22,	23,		24,	21,	20,	19,	18,	7.	Un .	<u>ت</u>	8	ω.	ς,	ω.	e 2,	2, 1	ÇT.	1,	10,	2, 1	12,	11,	16,	e 17,	Sequence 7, Appli	Today for nombro

ALIGNMENTS

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-09-452-565-1
Sequence 1, Application US/09452565
GENERAL INFORMATION:
APPLICANT: Anderson, Stephen F.
APPLICANT: Nash, Huw M.
APPLICANT: Felsch, Jason S.
TITLE OF INVENTION: ERYTHROPOIETIN RECEPTOR CHIMERA
FILE REFERENCE: 10845/011001
CURRENT APPLICATION NUMBER: US/09/452,565
CURRENT FILING DATE: 1999-12-01
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 676
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: -
OTHER INFORMATION: fusion protein including erythropoietin receptor extracellular
OTHER INFORMATION: domain, maltose binding proteins, linker, and GCN4 leucine
OTHER INFORMATION: zipper domain

1 KFESKAALLAARGPEELLCFTERLEDLVCFFEEAASAGVGPGNFSFSFQLEDEPWKLCRL 60

US-09-452-565-1

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US-09-502-984-5
Sequence 5. Application US/09502984
Sequence 5. Application US/09502984
GENERAL INFORMATION:
APPLICANT: Luo, Peizhi
TITLE OF INVENTION: STRUCTURE-BASED SCREENING TECHNIQUES
FITLE OF INVENTION NUMBER: US/09/502,984
CU37-ENT APPLICATION NUMBER: US/09/502,984
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GENERAL INFORMATION:
APPLICANT: Luc, Peishi
TITLE OF INVENTION: STRUCTURE-BASED SCREENING TECHNIQUES
FILE REFERENCE: A-68126-L/RFT/RMS/RMK
CURRENT APPLICARION UNBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
CURRENT FILING DATE: 2000-02-11
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 85.5%; Score 1098; DB 19; Best Local Similarity 100.0%; Pred. No. 2.1e-102; Matches 211; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 6
LENGTH: 211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/120,009
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
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TYPE: PRT
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                                               FOR DRUG DISCOVERY
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; SEQ ID NO 4
; LENGTH: 211
                                                                                                                                                     Matches
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CURRENT FILING DATE: 2000-02-1
PRIOR APPLICATION NUMBER: 60/1
PRIOR APPLICATION NUMBER: 60/1
PRIOR APPLICATION NUMBER: 60/1
PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 5
                                                                                                                                                                                                               ; FEATURE: ; OTHER INFORMATION: Description of Artificial Sequence: SYNTHETIC US-09-502-984-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Description
US-09-502-984-5
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                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: 60/120,009
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Luo, Peizhi
TITLE OF INVENTION: STRUCTURE-BASED SCREENING
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
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                                                           1 KFESKAALLAARGPEELLCFTERLEDLVCFFEEAASAGVGPGNFSFSFQLEDEPWKLCRL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 KFESKAALLAARGPEELLCFTERLEDLVCFWEEAASAGVGPGNYSFSYQLEDEPWKLCRL
PatentIn Ver. 2.1
                                                                                                                              Conservative
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97.28;
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Pred. No. 2.2e-100;
5; Mismatches 2;
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Pred. No. 1.
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                           NUMBER: PALE SOFTWARE: PALE SEQ ID NO 13
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SEQ ID NO 9
LENGTH: 211
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GENERAL INFORMATION:
APPLICANT: Luo, Peizhi
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Best Local
                                                                                                                             TITLE OF INVENTION: STRUCTURE-BASED SCRE
FILE REFERENCE: A-68126-1/RFT/RMS/RMS
CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: 60/120,009
PRIOR FILING DATE: 1999-02-11
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CURRENT FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: 60/120,009
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
                                                                                              PRIOR FILING DATE: 1999-04-29
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                                                                               NUMBER OF SEQ ID NOS:
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ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence
                  TYPE: PRT
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Pred. No. 4.5e
6; Mismatches
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SEQ ID NO 14
LENGTH: 211
TYPE: PRT
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Best Local Similarity
Matches 203; Conserv
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Best Local
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PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
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TITLE OF INVENTION: STRUCTURE-BASED SCREENING
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
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PRIOR FILING DATE: 1999-02-11
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                                                                      RLADESGHVVIRWLPPPETPMTSHIRFELDISAGNGAGSVQRVELLEGRTECVLSNLRGR 180
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TRYTFAVRARMAEPSFGGFWSAWSEPVSLLT
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Pred. No. 7.2e
6; Mismatches
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Pred. No. 7
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211
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7.2e-100;
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US-09-502-984-15

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RESULT 9
US-09-502-984-7
Sequence 7, Application US/09502984
GENERAL INFORMATION:
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SEQ ID NO 7
LENGTH: 211
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Best Local Similarity
Matches 202; Conserv
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GENERAL INFORMATION:
APPLICANT: Luo, Pelzhi
APPLICANT: Luo, Pelzhi
TITLE OF INVENTION: STRUCTURE-BASED SCREI
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
DEDICE APPLICATION WILLIAM
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TITLE OF INVENTION: STRUCTURE-BASED SCREENING TECHNIQUES FOR DRUG DISCOVERY
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
CURRENT FILING DATE: 2000-02-10
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Best Local Similarity
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PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/1
PRIOR FILING DATE: 1999-04-29
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NUMBER OF SEQ ID NOS:
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PRIOR TILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
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96.2%;
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95.7%;
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Pred. No. 7
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Pred. No. 1
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7.2e-100;
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SEQ ID NO 17
LENGTH: 211
                                                                                                                                              Sequence 16, Application US/09502984 GENERAL INFORMATION:
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Best Local Similarity
                                                                                   APPLICANT: Luo, Peizni
TITLE OF INVENTION: STRUCTURE-BASED SCREENING
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
                             CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: 60/1120,009
PRIOR FILING DATE: 1999-02-11
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PRIOR ETILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1999-04-29
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94.3%;
             60/131,674
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Pred. No. 1.5e-99;
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US-09-502-984-11
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Best Local S
Matches 198
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LENGTH: 211
TYPE: PRT
ORGANISM: Artificial Sequence
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SOFTWARE: Patentin V
SEQ ID NO 16
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                                                                                                                                                                                                                                                                           Query Match
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TITLE OF INVENTION: STRUCTURE-BASED SCREENING TECHNIQUES FOR DRUG DISCOVERY
EILE REFERENCE: A-68126-1/RFT/RMS/RMK
CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: 60/120,009
PRIOR APPLICATION NUMBER: 60/121,674
PRIOR PILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR APPLICATION NUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
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NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin V
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TYPE: PRT
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TRITIAVRARMAEPSFGGFWSAWSEPVSLLT 211
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                                                                                                                                                                                                                                       82.9%; Score 1064; DB 19; 94.3%; Pred. No. 5.9e-99; tive 10; Mismatches 2;
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Pred. No. 3.7e-99;
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; ORGANISM: Homo sapiens US-09-502-984-2
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US-09-502-984-2
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LENGTH: 211
TYPE: PRT
                                                                                                                                     SOFTWARE:
SEQ ID NO 2
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Query Match
Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: SCREENING TECHNIQUES FOR DRUG DISCOVERY
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/502,984
CURRENT FILING DATE: 2000-02-11
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TITLE OF INVENTION: ST
FILE REFERENCE: A-6812
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PRIOR EILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: 60/1
PRIOR FILING DATE: 1999-04-29
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                                                                                                   LENGTH: 211
TYPE: PRT
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94.3%;
82.6%;
93.8%;
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Pred. No. 1.2e
10; Mismatches
Score
Pred.
 1060; DB 19;
No. 1.5e-98;
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                  Length 211;
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Sequence 10, Application US/09502984

GENERAL INFORMATION:
APPLICANT: Luo, Peizhi
APPLICANT: Luo, Peizhi
TITLE OF INVENTION: STRUCTURE-BASED SCREENING TECHNIQUES FOR DRUG DISCOVERY
FILE REFERENCE: A-68126-1/RFT/RMS/RMK
CURRENT APPLICATION NUMBER: US/09/502,984

CURRENT APPLICATION NUMBER: 60/120,009
PRIOR APPLICATION NUMBER: 60/120,009
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION UNUMBER: 60/131,674
PRIOR FILING DATE: 1999-04-29
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
SEO ID NO 10
LENGTH: 211
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
Search completed: August 28, Job time: 515 sec
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                                                                                                                                                                                                                                                                                  Query Match 82.6%; Score 1060; DB 19; Best Local Similarity 93.8%; Pred. No. 1.5e-98; Matches 198; Conservative 11; Mismatches 2;
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                                                                                TRITIAVRARMAEPSFGGFWSAWSEPVSLLT 211
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